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5. The method according to claim 1, wherein said selected predetermined gDNA sequence within the 3'UTR or exon has a length of <u>at least</u> about 75 to about 2000 basesnucleotides.

Please add new claim 27, as follows:

27. The method according to claim 1, wherein said predetermined gDNA sequence within the 3'UTR or exon has a length of up to about 2000 nucleotides.